1/20

5

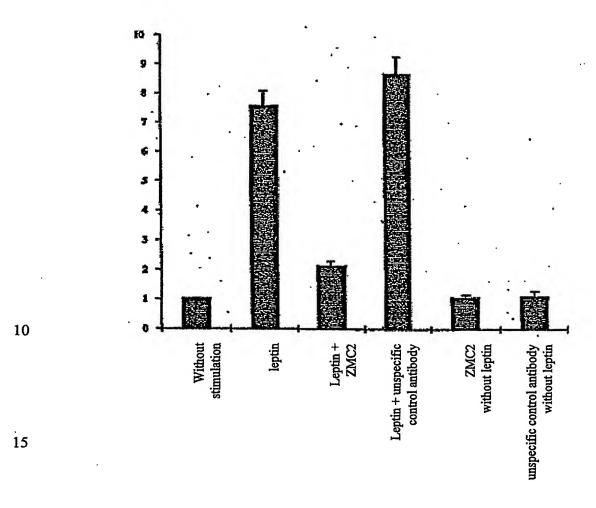


Figure 1

2/20

	A)	
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5		NVKSEDLADYFCLQHWNYPLTFGSGTKLEIKRADAAPTVSIFPPSSEQLT SGGASVVCFLNNFYPKDINVKWKIDGSERQNGVLNSWTDQDSKDSTYSMS STLTLTKDEYERHNSYTCEATHKTSTSPIVKSFNRGEC**SRVKRXQSXG
		GPGTPIRPIGXPYYNSLGGGFQ
	B)	
10		DNA: NANGTCATAATCCAATACCTATGCCTACGGCAGCCGCTGGATTGTTATTAC
		+3: X H N P I P M P T A A A G L L L L pComb3 vector SacI V _L (x) primer
1.5		DNA: TCGCTGCCCAACCAGCCATGGCCGAGCTCGTGATGACCCAGTCTCCAAAAT
15		+3: A A Q P A M A E L V M T Q S P K F
		DNA: TCATGTCCACATCAATAGGAGACAGGGTCAATATCACCTGCAAGGCCACTC +3: M S T S I G D R V N I T C K A T O
20		-
20	•	DNA: AGAATGTTCGTACTGCTGTTACCTGGTATCAACAGAAACCAGGGCAGTCTC . +3: N V R T A V T W Y Q Q K P G Q S P
	•	DNA: CTCAAGCACTGATTTCTTGGCATCCAACCGGCACACTGGTGTCCCTGCTC
25		+3: QALIFLASNRHTGVPAR
		DNA: GATTCACAGGCAGTGGATCTGGGACAGATTTCACTCTCACCATTAACAATG
	•	+3: FTGSGSGTDFTLTINNV
20		DNA: TGAAATCTGAAGACCTGGCAGATTATTTCTGTCTACAACATTGGAATTATC
30		+3: K S E D L A D Y F C L Q H W N Y P
		DNA: CTCTCACGTTCGGCTCGGGGACAAAGTTGGAAATAAAACGGGCTGATGCTG +3: L T F G S G T K L E I K R A D A A
. 35		DNA: CACCAACTGTATCCATCTTCCCACCATCCAGTGAGCAGTTAACATCTGGAG
		+3: PTVSIFPPSSEQLTSGG
		DNA: GTGCCTCAGTCGTGTGCTTCTTGAACAACTTCTACCCCAAAGACATCAATG
40		+3: A S V V C F L N N F Y P K D I N V
		DNA: TCAAGTGGAAGATTGATGGCAGTGAACGACAAAATGGCGTCCTGAACAGTT
		+3: K W K I D G S E R Q N G V L N S W
		Bcli DNA: GGACTGATCAGGACAGCAAAGACAGCACCTACAGCATGAGCAGCACCCTCA
45		+3: T D Q D S K D S T Y S M S S T L T
		DNA: CGTTGACCAAGGACGAGTATGAACGACATAACAGCTATACCTGTGAGGCCA
		+3: L T K D É Y E R H N S Y T C E A T
50		C _L (x) primer DNA: CTCACAAGACATCAACTTCACCCATTGTCAAGAGC <u>TTCAACAGGGGAGAGT</u>
50		+3: H K T S T S P I V K S F N R G E C
		Stop Xbal Notl Kpnl
		DNA: <u>GTTAGTAATCTAGA</u> GTTAA <u>GCGGCCGCAA</u> TCGAGGGGGGGCCC <u>GGTACC</u> CC +3: * * S R V K R P Q S R G G P V P Q
55		
		DNA: AATTCGCCCTATAGGGGNGCCGTATTACAATTCACTGGGCGGCGGTTTTCA
		+3: FAL * G X R I T I H W A A V F X
60		DNA: AN +3:

Figure 2

3/20

	A)																			
																				LESGP GSTYF
																				DYWGQ
5																				TVTWN
																				ASSTK
		VDK	KI	/PR	DC	TSF	НН	HHI	H*#	SL	VV	AVA	LH	SFV	XI	KAI	NRF	PA	K	•
	B)																			
10	-,	DNA:	тт	GGC	CNC	cce	ccc	TG6	ccc	cce	CA	דעע	יואיזי	ייתמי	יידיאור	ממי	יככי	CDC	AGTC	
		-1:	L	A	х	R	G	G	G	R	K	I	х	F	Х	R	E	T	V	
																			PACTC	
		-1:	I			vec			A	Y	G	P	A							
15		DNA:	GC	TGC	CCZ	LACC	AGO	CAI	:GGC	CCZ	GG:	'GAA	AC	rgc'i	hol 'CGA	. VH	D P	FILL	TGGC	
		-1:	A	A	Q	P	A	M	A	Q	V	K	L	L	E	s	G	P	G	
																			GTTC	
20	•		L																	
		DNA:	TC.	ATT	AAC	CGA	CGA	TGG	TGI	'AAG	CTC	GAT	TCG	GCA	'GCC	TCC	AGO	AAA	GGGT	
		-1:	s	и	Т	ע	ע	G	٧	S	W	1	R	Q	P	P	G	K	G.	
0.5		DNA:	CT	GGA	GTG	GCT	GGG	AGT	AAT	ATG	GGG	TGG	TGG	AAG	CAC	ATA	CTI	TAA	TTCA	
25			L																-	
		DNA:	CT'	TTT	CAA	ATC	CAG	ACT	GAG	CAT	CAC	CAG	GGA	CAA	CTC	TAA	GAG	CCA	AGTT	
		-1:	L	E.	K	S	R	ь	S	1	T	R	D	N	S	K	S	Q	V	
30		DNA:	TT	CTT.	AGA	AAT	GGA	CAG	TCT	ACA	AAC	TGA	TGA	CAC	AGC	CAT	GTA	CTA	CTGC	
		-1:	F	L	Е	M	D	s	L	Q	T	D	D	T	A	M	Y	Y	С	
		DNA:	GC	CAA	ACA	TGA	CGG	ACA	CGA	GAC	TAT	'GGA	CTA	TTG	GGG	TCA	AGG	AAC	CTCA	
35			A													-				
		DNA:	GTO	CAC	CGT	CTC	CTC	ATC	CAA	AAC	GAC	ACC	ccc	ATC	TGT	CTA	TCC	ACT	GCCC	
40		DNA:	P	G	S S	A A	TGC A	CCA	AAC T	TAA N	CTC	M	GGT V	GAC T	CCT T.	GGG	ATG	CCT t.	GGTC	
		DNA:	AAC	EGG(CTA	TTT	ccc	TGA	GCC	AGT	GAC	AGT	GAC	CTG	GAA	CTC	TGG	ATC	CCTG	
		-1:	K	G	I	r.	P	E	Þ	٧	T	V	Т	W	N	S	G	S	L	
45		DNA:	TCC	CAG	CGG	TGT	GCA	CAC	CTT	CCC	AGC	TGT	CCT	GCA	GTC	TGA	CCT	CTA	CACT	
		-1:	s	S	G	V	H	T	F	P	A	V	L	Q	s	D	L	Y	T	
		DNA:	CTC	AG	CAG	CTC	AGT	GAC'	TGT	ccc	CTC	CAG	CAC	CTG	GCC	CAG	CGA	GAC	CGTC	
50		-1:	L	S	S	S	V	T	V	P	S	S	T	W	P	S	E	T	V	
		DNA:	ACC	TG	CAA	CGT:	TGC	CCA	ccc	GGC	CAG	CAG	CAC	CAA	CCT	GG D	ממח	ממם	አ አጥጥ	
		-1:	T	С	N	V	A	Н	P	A	S	S	T	K	V	D	K	K	I	
55			C _H	1 (y	1) E	?rin	ner	Sp	eI		H	is t	ag			Sto	Œ			
		DNA:	<u>817</u>	P	R R	<u>عتع</u> ظر	C C	TAC.	S	H H	Η	TCA'	H H	rCA'	rca'	T'A	agc a	rage 2	CCTA T.	
		DNA:	GTG	GTO	GC	GGT(GGC'	TCT	CCA'	TTC	GTT'	TGT	GAN	SAT	AAA	GGC	CAA	TCG	NAGA	
		-1:	V	٧	A	٧	A	L	H	S	F'	۷	X	I	K	A	N	R	R	•
60		DNA:	CCI	GC1	ICN	A.														
		-1:	P	A	X															

Figure 3

4/20

	A) ·
	ATNCTTTNTTGTTCCTTTCTATGCGGCCCAGCCGGCCATGGCCCAGGTCCAGCTG
	CAGGAGTCAGGAACTGAAGTGGTAAAGCCTGGGGCTTCAGTGAAGTTGTCCT
	<u>GCAAGGCTTCTGGCTACATCTTCACAAGTTATGATATAGACTGGGTGAGGCAG</u>
5	<u>ACGCCTGA ACAGGGACTTGAGTGGATTGGATGGATTTTTCCTGGAGAGGGGA</u>
	<u>GTACTGAATACAATGAGAAGTTCAAGGGCAGGGCCACACTGAGTGTAGACAA</u>
	GTCCTCCA GCACAGCCTATATGGAGCTCACTAGGCTGACATCTGAGGACTCTG
	<u>CTGTCTATTTCTGTGCTAGAGGGGGACTACTATAGGCGCTACTTTGACTTGTGGG</u>
	<u>GCCAA</u> GGGACCACGGTCACCGTCTCCTCATGTGGAGGCGGTTCAGGCGGAGG
10	TGGCTCTGGCGGTGGCGGATCTGACATTGAGCTCACCCAGTCTCCAGCAATCA
	TGTCTGCATCTCCAGGGGAGAGGGTCACCATGACCTGCAGTGCCAGCTC
	AAGTATACGTTACATATTTGGTACCAACAGAAGCCTGGATCCTCCCCCA
	GACTCCTGATTTATGACACATCCAACGTGGCTCCTGGAGTCCCTTTTCGC
	TTCAGTGGCAGTGGGTCTGGGACCTCTTATTCTCTCACAATCAACCGAAT
15	GGAGGCTGAGGATGCTGCCACTTATTACTGCCAGGAGTGGAGTGGTTAT
	CCTCTCACGTTCGGCTCGGGCACCAAGCGGGAAATCAAACGGGCGGCCGC
	AGGTGCGCCGGTGCCGTATCCGGATCCGCTGGAACCGCGTGCCGCATAGACT- GTTGAA
	GIIGAA

20

B)

MAQVQLQESGTEVVKPGASVKLSCKASGYIFTSYDIDWVRQTPEQGLEWIG WIFPGEGSTEYNEKFKGRATLSVDKSSSTAYMELTRLTSEDSAVYFCARG DYYRRYFDLWGQGTTVTVSSGGGGSGGGGGGGGSDIELTQSPAIMSASP GERVTMTCSASSSIRYIYWYQQKPGSSPRLLIYDTSNVAPGVPFRFSGSG SGTSYSLTINRMEAEDAATYYCQEWSGYPLTFGSGTKREIKRAAAGAPVP YPDPLEPR

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5/20

A)

tcgctgcccaaccagccATGgcccaggtgaaactgctcgagtcaggacctggcctggtgg 5 cgccctcagagagcctgtccatcacatgcactatctcagggttctcattaaccgacgatg gtgtaagctggattcggcagcctccaggaaagggtctggagtggctgggagtaatatggg <u>gtggtggaagcacatactttaattcacttttcaaatccagactgagcatcaccagggaca</u> actctaagagccaagttttcttagaaatggacagtctacaaactgatgacacagccatgt actactgcgccaaacatgacggacacgagactatggactattggggtcaaggaacctcag 10 tcaccgtctcctcatccaaaacgacacccccatctgtctatccactggcccctggatctg ctgcccaaactaactccatggtgaccctgggatgcctggtcaagggctatttccctgagc cagtgacagtgacctggaactctggatccctgtccagcggtgtgcacaccttcccagctg tcctgcagtctgacctctacactctgagcagctcagtgactgtcccctccagcacctggc ccagcgagaccgtcacctgcaacgttgcccacccggccagcagcaccaaggtggacaaga 15 <u>aaattgtgcccagggattgtactagtggtggcggaggtagtggtggcggaggtagcggtg</u> <u>gcgqaggttctggtggcggaggttccgaattcctcgaggtgcccatccaaaaagtccaag</u> atgacaccaaaaccctcatcaagacaattgtcaccaggatcaatgacatttcacacacgc agtcagtctcctccaaacagaaagtcaccggtttggacttcattcctgggctccacccca tcctgaccttatccaagatggaccagacactggcagtctaccaacagatcctcaccagta 20 tgccttccagaaacgtgatccaaatatccaacgacctggagaacctccgggatcttcttc acgtgctggccttctctaagagctgccacttgccctgggccagtggcctggagaccttgg acagcctgqqqqqtqtcctgqaaqcttcagqctactccacagagqtqqtqqccctqaqca ggctgcaggggtctctgcaggacatgctgtggcagctggacctcagccctgggtgcacta gtcatcatcatcatcatcattaAgctagcctagtggtggcggtggctctcca 25

B)

40

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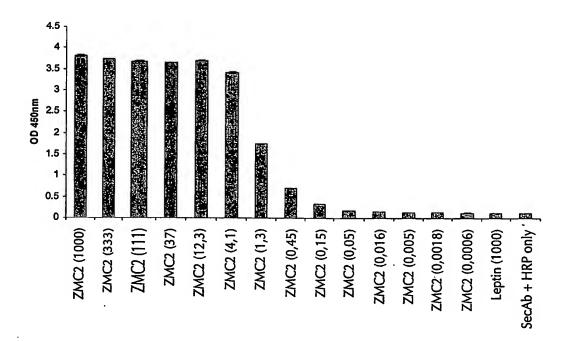


Figure 6

7/20

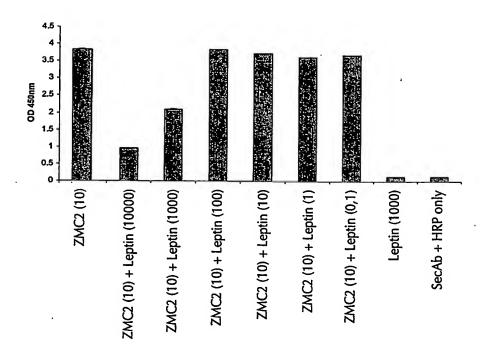


Figure 7

8/20

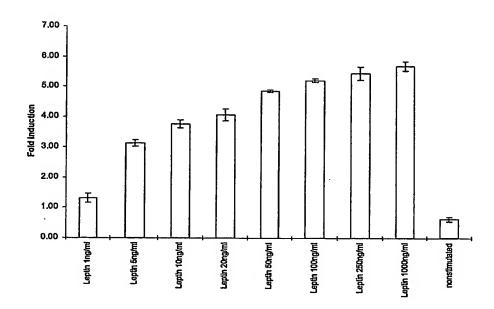


Figure 8

9/20

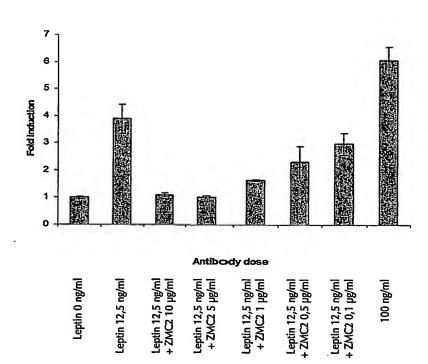


Figure 9

10/20

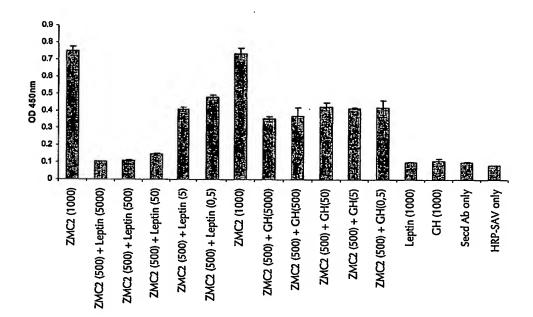


Figure 10

11/20

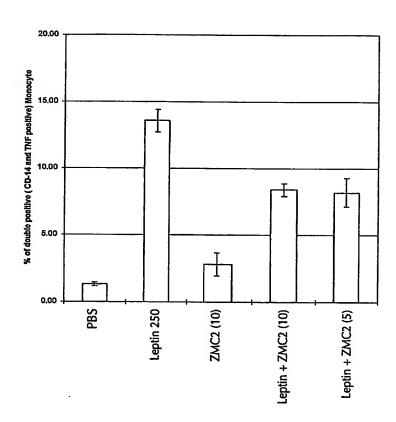
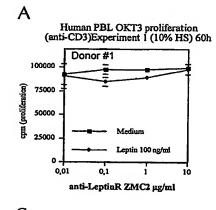
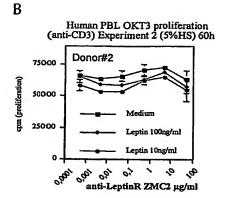


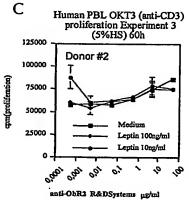
Figure 11

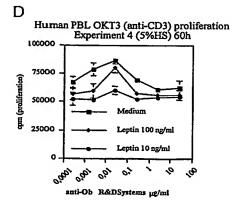
12/20

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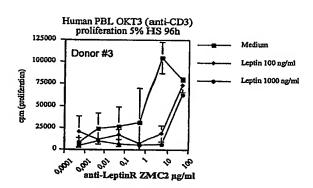
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Figure 12

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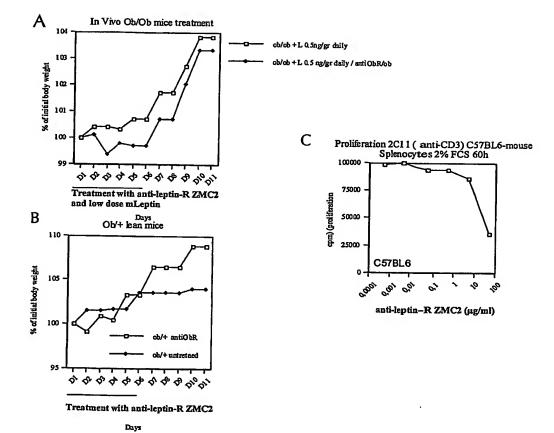


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Figure 13

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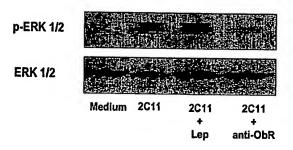
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Figure 14

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Figure 15

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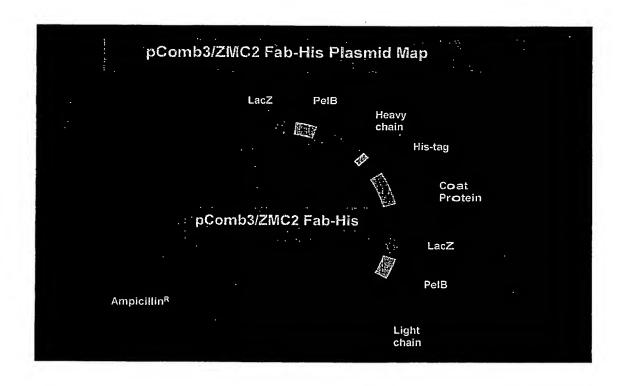


Figure 16

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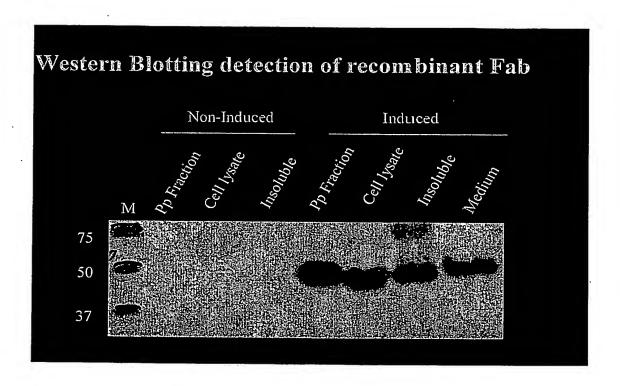


Figure 17

18/20

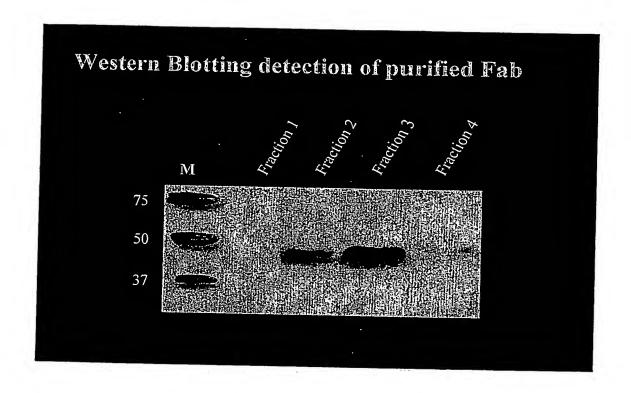


Figure 18

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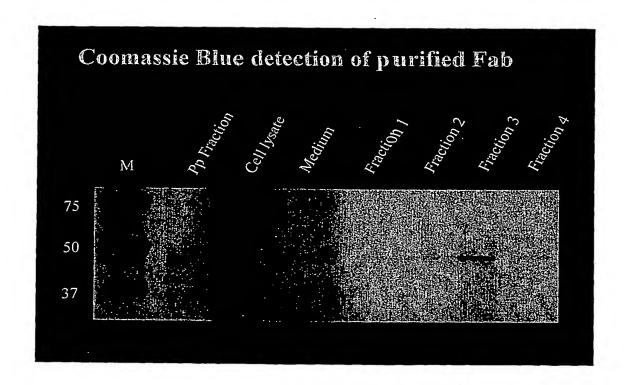
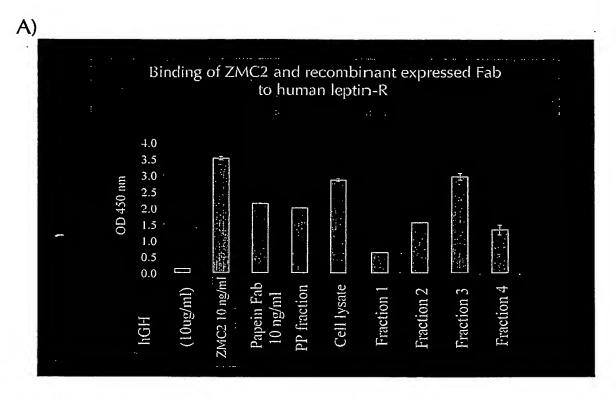


Figure 19

20/20



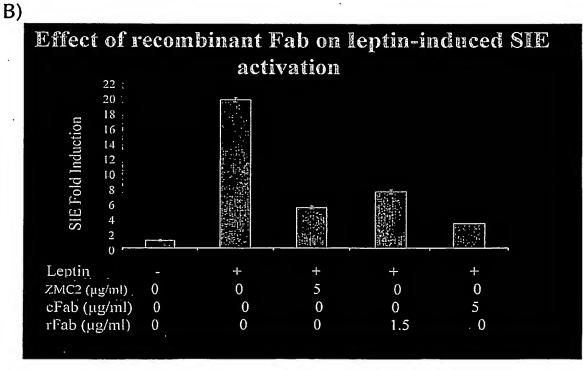


Figure 20